



Project 11

# SEQUENCE LISTING

<110> Gijzen, Mark

<120> Soybean Seed Coat Peroxidase Structural Gene And Regulatory Region

<130> 76-105

<140> US 08/939,905

<141> 1996-09-30

<150> US 08/723,414

<151> 1996-09-30

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<170> PatentIn version 3.0

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Tyr Arg Glu Thr Cys Pro Asn Leu Phe Pro Ile Val Phe Gly Val Ile  
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cgtcaggctt cactttcatg actgttttgt tctgggatgt gatgcctcag ttttgcctgaa	240
caatactgct acaatcgtaa gcgaacaaca agcttttcca aataacaact ctctaagggg	300
tttggatggt gtgaatcaga tcaaaactgc tgtagaaagt gcttgctcta acacagtttc	360
ttgtgctgat attcttgcac ttgctcaagc atcctctggt ctggcacaag gtcttagttg	420
gacggttcct ttaggaagaa gggatgggtt aaccgcaaac cgaacacttg caaatcaaaa	480
tcttcgggt ccattcaatt ccttggatca ccttaaactg catttgactg ctcaaggcct	540
cattactcct gttctagtgt cctctcggg tgctcatata tttggaagag ctcatcgcg	600
acaatttggt agtcgattgt acaacttcag cagtactgga agtcccgatc caactcttaa	660
cacaacttac ttacaacaac tgcgcacaat atgtcccaat ggtggacctg gcacaaacct	720
taccaatttc gatccaacga ctctgataa atttgacaag aactattact ccaatcttca	780
agtgaaaaag ggtttgctcc aaagtgatca agagtgttc tcaacttctg gtgcagatac	840
cattagcatt gtcgacaaat tcagcacoga tcaaaatgct ttctttgaga gctttaaggc	900
tgcaatgatt aaaatgggca atattggtgt gctaacaggg acaaaaggag agattagaaa	960
acaatgcaac tttgtgaact caaattctgc agaactagat ttagccacca tagcatccat	1020
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<210> 14

<211> 1200

<212> DNA

<213> Medicago sativa

<400> 14

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acccttttca aatgcacaac tagatccttc attttacaac agtacatggt ctaatcttga	120
ttcaatcgta cgtgggtgtgc tcacaaatgt ttcacaatct gatcccagaa tgcttggtag	180
tctcatcagg ctacattttc atgactgttt tgttcaagggt tgcgatgcct cgattttgct	240
gaacgatacg gctacaatag tgagcgagca aagtgcacca ccaaataaca actccataag	300

aggttttgat gtgataaacc agatcaaaac agcgggtggaa aatgcttggtc ctaacacagt 360  
 ttcttgtgct gatattcttg ctctttctgc tgaaatatca tctgatctgg caaatgggtcc 420  
 tacttggcaa gttccattag gaagaaggga tagtttgaca gcaaataatt ccttgcagc 480  
 tcaaaatctt cctgccccca ctttcaacct tactcgacta aaatctaact ttgataatca 540  
 aaacctcagt actactgac tagttgcact ctacaggtggc catacaattg gaagagggtca 600  
 atgcagatctt ttctgtgac gattatacaa tttcagcaac actggaaacc ccgattcaac 660  
 tcttaacacg acctatttac aaacattgca agcaatatgt cccaatgggtg gacctgggtac 720  
 aaacctaacg gatttggacc caaccacacc agatacatctt gactccaact actactccaa 780  
 tctccaagtt ggaaagggtc tgtttcagag tgaccaagag cttttttcca gaaatgggtc 840  
 tgacactatt tctattgtca atagtttcgc caataatcaa actctcttct ttgaaaattt 900  
 tgtagcctca atgataaaaa tgggtaatat tggagtttta actggatctc aaggtgaaat 960  
 tagaacacag tgtaatgctg tgaatgggaa ttcttctgga ttggctactg tagtcaccaa 1020  
 agaatcatca gaagatggaa tggctagctc attctaaata taagcttgga aaatattgaa 1080  
 gaggttctat aattttgtgc atacatatat ggtatgtgca tgtggtgtat tatgtttttg 1140  
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<210> 15

<211> 283

<212> PRT

<213> Glycine max

<400> 15

Phe	His	Asp	Cys	Phe	Val	Gln	Gly	Cys	Asp	Gly	Ser	Val	Leu	Leu	Asn
1				5				10						15	
Asn	Thr	Asp	Thr	Ile	Glu	Ser	Glu	Gln	Asp	Ala	Leu	Pro	Asn	Ile	Asn
			20				25						30		
Ser	Ile	Arg	Gly	Leu	Asp	Val	Val	Asn	Asp	Ile	Lys	Thr	Ala	Val	Glu
		35				40					45				
Asn	Ser	Cys	Pro	Asp	Thr	Val	Ser	Cys	Ala	Asp	Ile	Leu	Ala	Ile	Ala
	50					55				60					
Ala	Glu	Ile	Ala	Ser	Val	Ala	Gly	Arg	Arg	Ser	Gly	Trp	Pro	Val	Pro

65	70	75	80
Leu Gly Arg Arg Asp Ser Leu Thr Ala Asn Arg Thr Leu Ala Asn Gln			
	85	90	95
Asn Leu Pro Ala Pro Phe Phe Asn Leu Thr Gln Leu Lys Ala Ser Phe			
	100	105	110
Ala Val Gln Gly Leu Asn Thr Leu Asp Leu Val Thr Leu Ser Gly Gly			
	115	120	125
His Thr Ser Gly Arg Ala Arg Cys Ser Thr Phe Ile Asn Arg Leu Tyr			
	130	135	140
Asn Phe Ser Asn Thr Gly Leu Ile His Leu Asp Thr Thr Tyr Leu Glu			
	145	150	155
Val Leu Arg Ala Arg Cys Pro Gln Asn Ala Thr Gly Asp Asn Leu Thr			
	165	170	175
Asn Leu Asp Leu Ser Thr Pro Asp Gln Phe Asp Asn Arg Tyr Tyr Ser			
	180	185	190
Asn Leu Leu Gln Leu Asn Gly Leu Leu Gln Ser Asp Gln Glu Arg Phe			
	195	200	205
Ser Thr Pro Gly Ala Asp Thr Ile Pro Leu Ser Ile Ala Ser Ala Asn			
	210	215	220
Glu Asn Thr Phe Phe Ser Asn Phe Arg Val Ser Met Ile Lys Met Gly			
	225	230	235
Asn Ile Gly Val Leu Thr Gly Asp Glu Gly Glu Ile Arg Leu Gln Cys			
	245	250	255
Asn Phe Val Asn Gly Asp Ser Phe Gly Leu Ala Ser Val Ala Ser Lys			
	260	265	270
Asp Ala Lys Gln Lys Leu Val Ala Gln Ser Lys			
	275	280	

<210> 16

<211> 355

<212> PRT

<213> Medicago sativa

<400> 16

Met Asn Ser Leu Arg Ala Val Ala Ile Ala Leu Cys Cys Ile Val Val			
	1	5	10

Val Leu Gly Gly Leu Pro Phe Ser Ser Asn Ala Gln Leu Asp Pro Ser



20

25

30

Phe Tyr Arg Asn Thr Cys Pro Asn Val Ser Ser Ile Val Arg Glu Val  
35 40 45

Ile Arg Ser Val Ser Lys Lys Asp Pro Arg Met Leu Ala Ser Leu Val  
50 55 60

Arg Leu His Phe His Asp Cys Phe Val Gln Gly Cys Asp Ala Ser Val  
65 70 75 80

Leu Leu Asn Lys Thr Asp Thr Val Val Ser Glu Gln Asp Ala Phe Pro  
85 90 95

Asn Arg Asn Ser Leu Arg Gly Leu Asp Val Val Asn Gln Ile Lys Thr  
100 105 110

Ala Val Glu Lys Ala Cys Pro Asn Thr Val Ser Cys Ala Asp Ile Leu  
115 120 125

Ala Leu Ser Ala Glu Leu Ser Ser Thr Leu Ala Asp Gly Pro Asp Trp  
130 135 140

Lys Val Pro Leu Gly Arg Arg Asp Gly Leu Thr Ala Asn Gln Leu Leu  
145 150 155 160

Ala Asn Gln Asn Leu Pro Ala Pro Phe Asn Thr Thr Asp Gln Leu Lys  
165 170 175

Ala Ala Phe Ala Ala Gln Gly Leu Asp Thr Thr Asp Leu Val Ala Leu  
180 185 190

Ser Gly Ala His Thr Phe Gly Arg Ala His Cys Ser Leu Phe Val Ser  
195 200 205

Arg Leu Tyr Asn Phe Ser Gly Thr Gly Ser Pro Asp Pro Thr Leu Asn  
210 215 220

Thr Thr Tyr Leu Gln Gln Leu Arg Thr Ile Cys Pro Asn Gly Gly Pro  
225 230 235 240

Gly Thr Asn Leu Thr Asn Phe Asp Pro Thr Thr Pro Asp Lys Phe Asp  
245 250 255

Lys Asn Tyr Tyr Ser Asn Leu Gln Val Lys Lys Gly Leu Leu Gln Ser  
260 265 270

Asp Gln Glu Leu Phe Ser Thr Ser Gly Ser Asp Thr Ile Ser Ile Val  
275 280 285

Asn Lys Phe Ala Thr Asp Gln Lys Ala Phe Phe Glu Ser Phe Arg Ala  
290 295 300

Ala Met Ile Lys Met Gly Asn Ile Gly Val Leu Thr Gly Asn Gln Gly  
305 310 315 320

Glu Ile Arg Lys Gln Cys Asn Phe Val Asn Ser Lys Ser Ala Glu Leu

325

330

335

Gly Leu Ile Asn Val Ala Ser Ala Asp Ser Ser Glu Glu Gly Met Val  
 340 345 350

Ser Ser Met  
 355

<210> 17

<211> 358

<212> PRT

<213> Medicago sativa

<400> 17

Met Asn Ser Leu Ala Thr Ser Met Trp Cys Val Val Leu Leu Val Val  
 1 5 10 15

Leu Gly Gly Leu Pro Phe Ser Ser Asp Ala Gln Leu Ser Pro Thr Phe  
 20 25 30

Tyr Ser Lys Thr Cys Pro Thr Val Ser Ser Ile Val Ser Asn Val Leu  
 35 40 45

Thr Asn Val Ser Lys Thr Asp Pro Arg Met Leu Ala Ser Leu Val Arg  
 50 55 60

Leu His Phe His Asp Cys Phe Val Leu Gly Cys Asp Ala Ser Val Leu  
 65 70 75 80

Leu Asn Asn Thr Ala Thr Ile Val Ser Glu Gln Gln Ala Phe Pro Asn  
 85 90 95

Asn Asn Ser Leu Arg Gly Leu Asp Val Val Asn Gln Ile Lys Leu Ala  
 100 105 110

Val Glu Val Pro Cys Pro Asn Thr Val Ser Cys Ala Asp Ile Leu Ala  
 115 120 125

Leu Ala Ala Gln Ala Ser Ser Val Leu Ala Gln Gly Pro Ser Trp Thr  
 130 135 140

Val Pro Leu Gly Arg Arg Asp Gly Leu Thr Ala Asn Arg Thr Leu Ala  
 145 150 155 160

Asn Gln Asn Leu Pro Ala Pro Phe Asn Ser Leu Asp Gln Leu Lys Ala  
 165 170 175

Ala Phe Thr Ala Gln Gly Leu Asn Thr Thr Asp Leu Val Ala Leu Ser  
 180 185 190

Gly Ala His Thr Phe Gly Arg Ala His Cys Ala Gln Phe Val Ser Arg

195                      200                      205  
 Leu Tyr Asn Phe Ser Ser Thr Gly Ser Pro Asp Pro Thr Leu Asn Thr  
     210                      215                      220  
 Thr Tyr Leu Gln Gln Leu Arg Thr Ile Cys Pro Asn Gly Gly Pro Gly  
     225                      230                      235                      240  
 Thr Asn Leu Thr Asn Phe Asp Pro Thr Thr Pro Asp Lys Phe Asp Lys  
     245                      250                      255  
 Asn Tyr Tyr Ser Asn Leu Gln Val Lys Lys Gly Leu Leu Gln Ser Asp  
     260                      265                      270  
 Gln Glu Leu Phe Ser Thr Ser Gly Ala Asp Thr Ile Ser Ile Val Asn  
     275                      280                      285  
 Lys Phe Ser Thr Asp Gln Asn Ala Phe Phe Glu Ser Phe Lys Ala Ala  
     290                      295                      300  
 Met Ile Lys Met Gly Asn Ile Gly Val Leu Thr Gly Thr Lys Gly Glu  
     305                      310                      315                      320  
 Ile Arg Lys Gln Cys Asn Phe Val Asn Phe Val Asn Ser Asn Ser Ala  
     325                      330                      335  
 Glu Leu Asp Leu Ala Thr Ile Ala Ser Ile Val Glu Ser Leu Glu Asp  
     340                      345                      350  
 Gly Ile Ala Ser Val Ile  
     355

<210> 18

<211> 347

<212> PRT

<213> Medicago sativa

<400> 18

Met Trp Cys Val Val Leu Leu Val Val Leu Gly Gly Leu Pro Phe Ser  
     1                      5                      10                      15  
 Ser Asp Ala Gln Leu Ser Pro Thr Phe Tyr Ser Lys Thr Cys Pro Thr  
     20                      25                      30  
 Val Ser Ser Ile Val Ser Asn Val Leu Thr Asn Val Ser Lys Thr Asp  
     35                      40                      45  
 Pro Arg Met Leu Ala Ser Leu Val Arg Leu His Phe His Asp Cys Phe  
     50                      55                      60  
 Val Leu Gly Cys Asp Ala Ser Val Leu Leu Asn Asn Thr Ala Thr Ile

65	70	75	80
Val Ser Glu Gln Gln Ala Phe Pro Asn Asn Asn Ser Leu Arg Gly Leu			
	85	90	95
Asp Val Val Asn Gln Ile Lys Thr Ala Val Glu Ser Ala Cys Pro Asn			
	100	105	110
Thr Val Ser Cys Ala Asp Ile Leu Ala Leu Ala Gln Ala Ser Ser Val			
	115	120	125
Leu Ala Gln Gly Pro Ser Trp Thr Val Pro Leu Gly Arg Arg Asp Gly			
	130	135	140
Leu Thr Ala Asn Arg Thr Leu Ala Asn Gln Asn Leu Pro Ala Pro Phe			
	145	150	155
Asn Ser Leu Asp His Leu Lys Leu His Leu Thr Ala Gln Gly Leu Ile			
	165	170	175
Thr Pro Val Leu Val Ala Leu Ser Gly Ala His Thr Phe Gly Arg Ala			
	180	185	190
His Cys Ala Gln Phe Val Ser Arg Leu Tyr Asn Phe Ser Ser Thr Gly			
	195	200	205
Ser Pro Asp Pro Thr Leu Asn Thr Thr Tyr Leu Gln Gln Leu Arg Thr			
	210	215	220
Ile Cys Pro Asn Gly Gly Pro Gly Thr Asn Leu Thr Asn Phe Asp Pro			
	225	230	235
Thr Thr Pro Asp Lys Phe Asp Lys Asn Tyr Tyr Ser Asn Leu Gln Val			
	245	250	255
Lys Lys Gly Leu Leu Gln Ser Asp Gln Glu Leu Phe Ser Thr Ser Gly			
	260	265	270
Ala Asp Thr Ile Ser Ile Val Asp Lys Phe Ser Thr Asp Gln Asn Ala			
	275	280	285
Phe Phe Glu Ser Phe Lys Ala Ala Met Ile Lys Met Gly Asn Ile Gly			
	290	295	300
Val Leu Thr Gly Thr Lys Gly Glu Ile Arg Lys Gln Cys Asn Phe Val			
	305	310	315
Asn Ser Asn Ser Ala Glu Leu Asp Leu Ala Thr Ile Ala Ser Ile Val			
	325	330	335
Glu Ser Leu Glu Asp Gly Ile Ala Ser Val Ile			
	340	345	

<210> 19

<211> 351

<212> PRT

<213> Medicago sativa

<400> 19

Met Leu Gly Leu Ser Ala Thr Ala Phe Cys Cys Met Val Phe Val Leu  
1 5 10 15

Ile Gly Gly Val Pro Phe Ser Asn Ala Gln Leu Asp Pro Ser Phe Tyr  
20 25 30

Asn Ser Thr Cys Ser Asn Leu Asp Ser Ile Val Arg Gly Val Leu Thr  
35 40 45

Asn Val Ser Gln Ser Asp Pro Arg Met Leu Gly Ser Leu Ile Arg Leu  
50 55 60

His Phe His Asp Cys Phe Val Gln Gly Cys Asp Ala Ser Ile Leu Leu  
65 70 75 80

Asn Asp Thr Ala Thr Ile Val Ser Glu Gln Ser Ala Pro Pro Asn Asn  
85 90 95

Asn Ser Ile Arg Gly Leu Asp Val Ile Asn Gln Ile Lys Thr Ala Val  
100 105 110

Glu Asn Ala Cys Pro Asn Thr Val Ser Cys Ala Asp Ile Leu Ala Leu  
115 120 125

Ser Ala Glu Ile Ser Ser Asp Leu Ala Asn Gly Pro Thr Trp Gln Val  
130 135 140

Pro Leu Gly Arg Arg Asp Ser Leu Thr Ala Asn Asn Ser Leu Ala Ala  
145 150 155 160

Gln Asn Leu Pro Ala Pro Thr Phe Asn Leu Thr Arg Leu Lys Ser Asn  
165 170 175

Phe Asp Asn Gln Asn Leu Ser Thr Thr Asp Leu Val Ala Leu Ser Gly  
180 185 190

Gly His Thr Ile Gly Arg Gly Gln Cys Arg Phe Phe Val Asp Arg Leu  
195 200 205

Tyr Asn Phe Ser Asn Thr Gly Asn Pro Asp Ser Thr Leu Asn Thr Thr  
210 215 220

Tyr Leu Gln Thr Leu Gln Ala Ile Cys Pro Asn Gly Gly Pro Gly Thr  
225 230 235 240

Asn Leu Thr Asp Leu Asp Pro Thr Thr Pro Asp Thr Phe Asp Ser Asn  
245 250 255

Tyr Tyr Ser Asn Leu Gln Val Gly Lys Gly Leu Phe Gln Ser Asp Gln

260	265	270
Glu Leu Phe Ser Arg Asn Gly Ser Asp Thr Ile Ser Ile Val Asn Ser		
275	280	285
Phe Ala Asn Asn Gln Thr Leu Phe Phe Glu Asn Phe Val Ala Ser Met		
290	295	300
Ile Lys Met Gly Asn Ile Gly Val Leu Thr Gly Ser Gln Gly Glu Ile		
305	310	315
Arg Thr Gln Cys Asn Ala Val Asn Gly Asn Ser Ser Gly Leu Ala Thr		
325	330	335
Val Val Thr Lys Glu Ser Ser Glu Asp Gly Met Ala Ser Ser Phe		
340	345	350

<210> 20

<211> 22

<212> DNA

<213> Medicago sativa

<400> 20

taaaatcata tcagcttact cc